

BMS 353 Bioinformatics for Biomedical Science

Module coordinator: Dr Marta Milo Research Software Instructor: Dr Mike Croucher



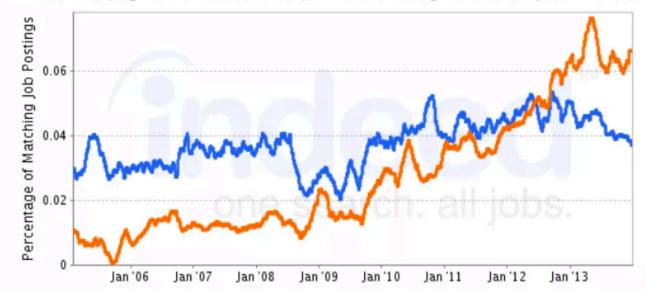




R is a free language and environment for statistical computing and graphics.

Job Trends from Indeed.com

R !"R D" !"A R" !"H R" !"R N" !toys !kids !" R Walgreen" !walmart !"HVAC R" !"R Bard" and (
 SPSS and ("big data" or "statistical analysis" or "data mining" or "data analytics" or "machir









Packages Save time by using other people's code.

Comprehensive R Archive Network (CRAN) 7468 packages (November 2015)

https://cran.r-project.org/web/packages/





Bioconductor 1104 packages (November 2015)

https://www.bioconductor.org/





Packages Save time by using other people's code.

Comprehensive R Archive Network (CRAN) 9262 packages (October 2016)

https://cran.r-project.org/web/packages/





Bioconductor 1211 packages (October 2016)

https://www.bioconductor.org/





Open source, interactive data science and scientific computing across over 40 programming languages.







Currently in use at







Bloomberg









































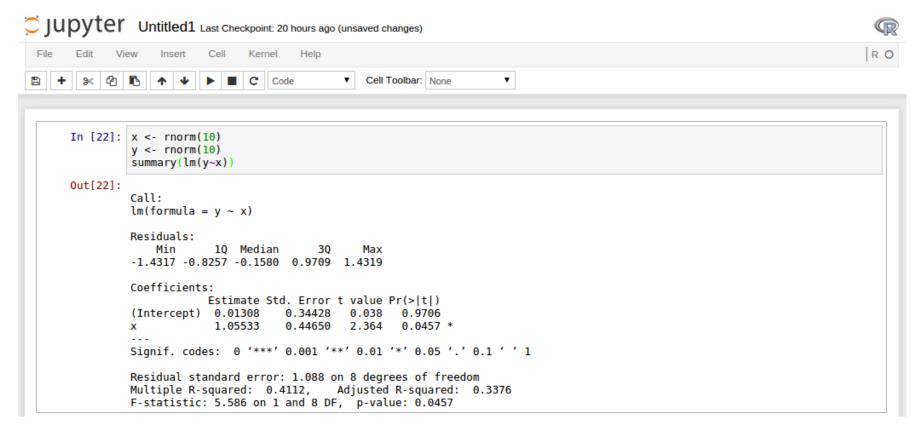
Combine live computer code, data, text and mathematics in one interactive document.

Academic papers only give a description of your analysis.

Jupyter notebooks contain both the description and the analysis itself.













Collaborative computational mathematics
An open source environment for running Jupyter notebooks (and others!) in the cloud.

Dedicated Virtual Machine for this course on the Google Cloud Platform.

As powerful as a node on the Sheffield University
Supercomputer: Iceberg

THE 1ST





All you need is a web browser and internet connection.





Log into SageMathCloud https://cloud.sagemath.com/

Create account (or sign in)
☐ First, agree to the Terms of Service
Use your email address
Name
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g Google Facebook Github Twitter
Not working? Email us at help@sagemath.com immediately!

